

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/935,799

DATE: 11/08/2001

TIME: 13:22:18

Input Set : A:\es.txt

Output Set: N:\CRF3\11082001\I935799.raw

1 <110> APPLICANT: MOCKEL, Bettina <120>
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W--> 3 <140> CURRENT APPLICATION NUMBER: US/09/935,799
W--> 0 <160> NUMBER OF SEQ ID NOS:
3 <170> SOFTWARE: PatentIn version 3.1<210> 1<211> 2718<212> DNA<213> Corynebacterium

Does Not Comply
Corrected Diskette Needed

ERRORED SEQUENCES

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*major format
error (see
next page)*

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11 tggccgaact ttcccttttc tgcattgcatt tctgcacaca gtttctgccc gctgtttctg      180
13 cccgctgttt ctacgcata gtg gct ttg aaa cga ccc gaa gag aaa aca gta      232
14                               Met Ala Leu Lys Arg Pro Glu Glu Lys Thr Val
15                               1               5               10
17 aag atc gtg acc ata aaa cag act gac aac atc aat gac gat gat ttg      280
18 Lys Ile Val Thr Ile Lys Gln Thr Asp Asn Ile Asn Asp Asp Asp Leu
19                               15               20               25
21 gtg tac agc aac gct act gac ctt cca gta ggc gtg aag aag tcc cct      328
22 Val Tyr Ser Asn Ala Thr Asp Leu Pro Val Gly Val Lys Lys Ser Pro
23                               30               35               40
25 aaa atg tca ccg acc gcc cgc gtt ggt ctc ctt gtc ttt ggg gtt atc      376
26 Lys Met Ser Pro Thr Ala Arg Val Gly Leu Leu Val Phe Gly Val Ile
27                               45               50               55
29 gcg gcg gtg ggt tgg gga gca atc gct ttc tcc cgt ggc gaa aca atc      424
30 Ala Ala Val Gly Trp Gly Ala Ile Ala Phe Ser Arg Gly Glu Thr Ile
31 60                               65               70               75
33 aac tct gtg tgg ctg gtt ttg gcg gca gtt ggt tcc tat atc att gcg      472
34 Asn Ser Val Trp Leu Val Leu Ala Ala Val Gly Ser Tyr Ile Ile Ala
35                               80               85               90
37 ttt tct ttc tat gcc cga ctg att gaa tac aaa gtt gtt aag ccg aaa      520
38 Phe Ser Phe Tyr Ala Arg Leu Ile Glu Tyr Lys Val Val Lys Pro Lys
39                               95               100              105
41 gat cag cga gca acc ccg gcg gaa tac gtt aat gac ggc aag gac tat      568
42 Asp Gln Arg Ala Thr Pro Ala Glu Tyr Val Asn Asp Gly Lys Asp Tyr
43                               110              115              120
45 gtc cca acg gat cgt cgt gtg ctt ttt ggc cac cac ttt gca gct att      616
46 Val Pro Thr Asp Arg Arg Val Leu Phe Gly His His Phe Ala Ala Ile
47                               125              130              135

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